STIC Biotechnology Systems Branch

RAW SEQUENCE LISTING ERROR REPORT



The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number:	10/562,324
Source:	JFWP 01/09/2006
Date Processed by STIC:	01/09/2006

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.
PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 4.4.0 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

- 1. EFS-Bio (http://www.uspto.gov/ebc/efs/downloads/documents.htm, EFS Submission User Manual ePAVE)
- 2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
- 3. Hand Carry, Federal Express, United Parcel Service, or other delivery service (EFFECTIVE 01/14/05): U.S. Patent and Trademark Office, Mail Stop Sequence, Customer Window, Randolph Building, 401 Dulany Street, Alexandria, VA 22314

Revised 01/10/06





IFWO

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/562,324

DATE: 01/09/2006

TIME: 10:43:50

Input Set : A:\PTO.DA.txt

Output Set: N:\CRF4\01092006\J562324.raw

5 <110> APPLICANT: Bayer AG

9 <120> TITLE OF INVENTION: NEUES EIMERIA GEN UND PROTEIN SOWIE DEREN

VERWENDUNG

13 <130> FILE REFERENCE: LeA 36695 DE

C--> 17 <140> CURRENT APPLICATION NUMBER: US/10/562,324

C--> 17 <141> CURRENT FILING DATE: 2005-12-23

17 <160> NUMBER OF SEQ ID NOS: 24

21 <170> SOFTWARE: PatentIn version 3.1

ERRORED SEQUENCES

563 <210> SEQ ID NO: 24

565 <211> LENGTH: 18

567 <212> TYPE: DNA

569 <213> ORGANISM: Artificial Sequence

573 <220> FEATURE:

575 <223> OTHER INFORMATION: Primer BGH reverse

) Pls delete

577 <400> SEQUENCE: 24

578 tagaaggcac agtcgagg

R--> 584 Le a 36 695

E--> 587\- 11 -

Does Not Comply Corrected Diskette Needed

18

VERIFICATION SUMMARY

DATE: 01/09/2006

PATENT APPLICATION: US/10/562,324

TIME: 10:43:51

Input Set : A:\PTO.DA.txt

Output Set: N:\CRF4\01092006\J562324.raw

L:17 M:270 C: Current Application Number differs, Replaced Current Application No

L:17 M:271 C: Current Filing Date differs, Replaced Current Filing Date L:45 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:1,Line#:41 L:584 M:254 E: No. of Bases conflict, LENGTH:Input:695 Counted:21 SEQ:24 L:584 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:2

L:584 M:112 C: (48) String data converted to lower case,

L:587 M:254 E: No. of Bases conflict, this line has no nucleotides.

L:587 M:252 E: No. of Seq. differs, <211> LENGTH:Input:18 Found:21 SEQ:24

VERIFICATION SUMMARY

DATE: 01/09/2006

PATENT APPLICATION: US/10/562,324

TIME: 10:43:51

Input Set : A:\PTO.DA.txt

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L:17 M:270 C: Current Application Number differs, Replaced Current Application No

L:17 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:45 M:258 W: Mandatory Peature missing, <223> Blank for SEQ#:1, Line#:41

L:584 M:254 E: No. of Bases conflict, LENGTH:Input:695 Counted:21 SEQ:24 L:584 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:2

L:584 M:112 C: (48) String data converted to lower case,

L:587 M:254 B: No. of Bases conflict, this line has no nucleotides.

L:587 M:252 B: No. of Seq. differs, <211> LENGTH:Input:18 Found:21 SEQ:24